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## Amendments to the Claims

This listing of claims will replace all prior versions and listings of claims in the application.

- 1. (Previously Presented) An isolated nucleic acid molecule encoding an odorant receptor protein, wherein the receptor protein comprises seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, and is further characterized by at least one of the following characteristics:
  - (a) the loop between the first transmembrane domain and the second transmembrane domain, and the second transmembrane domain together comprise consecutive amino acids having the following sequence: -L, X, X, P, M, Y, X, F, L- (SEQ ID NO: 55);
  - (b) the third transmembrane domain, and the loop between the third transmembrane domain and the fourth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-M, X, Y, D, R, X, X, A, I, C- (SEQ ID NO: 57); or

-D, R, X, X, A, I, C- (SEQ ID NO: 59);

(c) the loop between the fifth transmembrane domain and the sixth transmembrane domain, and the sixth transmembrane domain together comprise consecutive amino acids having

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one of the following sequences:

-K or R, X, F, S, T, C, X, S, H- (SEQ ID NO: 61); or

-F, S, T, C, X, S, H- (SEQ ID NO: 63); or

(d) the seventh transmembrane domain and the C-terminal domain together comprise consecutive amino acids having one of the following sequences:

-P, X, X, N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 65); or

-P, X, X, N, P, X, I, Y- (SEQ ID NO: 67); or

-N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 69);

wherein X is any amino acid.

- 2. (Original) The isolated nucleic acid molecule of claim 1 wherein:
  - (a) the loop between the first transmembrane domain and the second transmembrane domain, and the second transmembrane domain together comprise consecutive amino acids having the following sequence:

-L, H or Q, K or M or T, PMY, F or L, FL- (SEQ ID NO: 56);

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(b) the third transmembrane domain, and the loop between the third transmembrane domain and the fourth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-M, A or S, YDR, F or Y, L or V, AIC- (SEQ ID NO: 58); or

-DR, F or Y, L or V, AIC- (SEQ ID NO: 60);

(c) the loop between the fifth transmembrane domain and the sixth transmembrane domain, and the sixth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-K or R, A or I or S or V, FSTC, A or G or S, SH- (SEQ ID NO: 62); or

-FSTC, A or G or S, SH- (SEQ ID NO: 64); or

(d) the seventh transmembrane domain and the C-terminal domain together comprise consecutive amino acids having one of the following sequences:

-P, M or L or V, F or L or V, NP, F or I, IY, C or S or T, LRN- (SEQ ID NO: 66); or

 $-\dot{P}$ , M or L or V, F or L or V, NP, F or I, IY- (SEQ ID NO: 68); or

-NP, F or I, IY, C or S or T, LRN- (SEQ ID NO: 70).

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3. (Original) The isolated nucleic acid molecule of claim 1, wherein the receptor protein is characterized by at least two of the characteristics (a) through (d).

- 4. (Original) The isolated nucleic acid molecule of claim 1, wherein the receptor protein is characterized by at least three of the characteristics (a) through (d).
- 5. (Original) The isolated nucleic acid molecule of claim 1, wherein the receptor protein is characterized by all of the characteristics (a) through (d).
- 6. (Previously Presented) An isolated nucleic acid molecule encoding an odorant receptor protein comprising seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, wherein the nucleic acid molecule encodes a protein selected from the group consisting of:
  - (a) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with tyrosine at position 333 as set forth in row F3 of Figures 4A to 4M (SEQ ID NO: 71),
  - (b) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with glutamine at position 313 as set forth in row F5 of Figures 4A to 4L

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(SEQ ID NO: 72),

- (c) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with lysine at position 311 as set forth in row F6 of Figures 4A to 4L (SEQ ID NO: 73),
- (d) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with glycine at position 317 as set forth in row F12 of Figures 4A to 4L (SEQ ID NO: 74),
- (e) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 310 as set forth in row I3 of Figures 4A to 4L (SEQ ID NO: 75),
- (f) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with glycine at position 327 as set forth in row I7 of Figures 4A to 4L (SEQ ID NO: 76),
- (g) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with tryptophan at position 312 as set forth in row I8 of Figures 4A to 4L

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(SEQ ID NO: 77),

- (h) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 314 as set forth in row I9 of Figures 4A to 4L (SEQ ID NO: 78),
- (i) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 312 as set forth in row I14 of Figures 4A to 4L (SEQ ID NO: 79),
- (j) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 314 as set forth in row I15 of Figures 4A to 4L (SEQ ID NO: 80); and
- (k) an odorant receptor protein that shares from 40-80% amino acid identity with any one of the proteins of (a)-(j), comprises seven transmembrane domains, and is further characterized by at least one of the following characteristics:
  - (i) the loop between the first transmembrane domain and the second transmembrane domain, and the second transmembrane domain together comprise consecutive amino acids having the following sequence: -L, X,

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X, P, M, Y, X, F, L- (SEQ ID NO: 55);

(ii) the third transmembrane domain, and the loop between the third transmembrane domain and the fourth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-M, X, Y, D, R, X, X, A, I, C- (SEQ ID NO: 57); or

-D, R, X, X, A, I, C- (SEQ ID NO: 59);

(iii) the loop between the fifth transmembrane domain and the sixth transmembrane domain, and the sixth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-K or R, X, F, S, T, C, X, S, H- (SEQ ID NO: 61); or

-F, S, T, C, X, S, H- (SEQ ID NO: 63); or

(iv) the seventh transmembrane domain and the C-terminal domain together comprise consecutive amino acids having one of the following sequences:

-P, X, X, N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 65); or

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-P, X, X, N, P, X, I, Y- (SEQ ID NO: 67); or -N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 69); wherein X is any amino acid.

- 7. (Original) The isolated nucleic acid molecule of claim 6 wherein:
  - (i) the loop between the first transmembrane domain and the second transmembrane domain, and the second transmembrane domain together comprise consecutive amino acids having the following sequence:
    - -L, H or Q, K or M or T, PMY, F or L, FL- (SEQ ID NO: 56);
  - (ii) the third transmembrane domain, and the loop between the third transmembrane domain and the fourth transmembrane domain together comprise consecutive amino acids having one of the following sequences:
    - -M, A or S, YDR, F or Y, L or V, AIC- (SEQ ID NO: 58); or
    - -DR, F or Y, L or V, AIC- (SEQ ID NO: 60);
  - (iii) the loop between the fifth transmembrane domain and the sixth transmembrane domain, and the sixth transmembrane domain together comprise consecutive amino acids having

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one of the following sequences:

-K or R, A or I or S or V, FSTC, A or G or S, SH- (SEQ ID NO: 62); or

-FSTC, A or G or S, SH- (SEQ ID NO: 64); or

- (iv) the seventh transmembrane domain and the .C-terminal domain together comprise consecutive amino acids having one of the following sequences:
  - -P, M or L or V, F or L or V, NP, F or I, IY, C or S or T, LRN- (SEQ ID NO: 66); or
  - -P, M or L or V, F or L or V, NP, F or I, IY- (SEQ ID NO: 68); or
  - -NP, F or I, IY, C or S or T, LRN- (SEQ ID NO: 70).
- 8. (Previously Presented) An isolated nucleic acid molecule encoding an odorant receptor protein comprising seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, wherein the nucleic acid molecule comprises a nucleic acid sequence which can be amplified by polymerase chain reaction using:
  - (a) any one of 5' primers A1 (SEQ ID NO: 37), A2 (SEQ ID NO: 38), A3 (SEQ ID NO: 39), A4 (SEQ ID NO: 40), or A5 (SEQ

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ID NO: 41); and

(b) any one of 3' primers B1 (SEQ ID NO: 42), B2 (SEQ ID NO: 43), B3 (SEQ ID NO: 44), B4 (SEQ ID NO: 45), B5 (SEQ ID NO: 46), or B6 (SEQ ID NO: 47).

## 9-12. (Canceled)

- 13. (Original) The isolated nucleic acid molecule of any one of claims 1, 6, 8, or 9, wherein the odorant receptor is a vertebrate odorant receptor.
- 14. (Original) The isolated nucleic acid molecule of claim 13, wherein the vertebrate odorant receptor is a fish odorant receptor or a mammalian odorant receptor.
- 15. (Original) The isolated nucleic acid molecule of claim 14, wherein the mammalian odorant receptor is a human odorant receptor, a rat odorant receptor, a mouse odorant receptor or a dog odorant receptor.
- 16. (Original) The isolated nucleic acid molecule of claim 1, 6,8, or 9, wherein the nucleic acid is DNA.
- 17. (Original) The isolated nucleic acid molecule of claim 16, wherein the DNA is cDNA.
- 18. (Original) A vector comprising the isolated nucleic acid

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molecule of claim 1, 6, 8, or 9.

19. (Original) The vector of claim 18, wherein the vector additionally comprises elements necessary for replication and expression in a suitable host.

- 20. (Canceled)
- 21. (Original) A cell transfected with the vector of claim 19.
- 22. (Original) The cell of claim 21, wherein the cell is an olfactory cell.
- 23. (Original) The cell of claim 21, wherein the cell is a nonolfactory cell.
- 24. (Original) The cell of claim 23, wherein prior to being transfected with the vector the non-olfactory cell does not express an odorant receptor protein.

## 25-63. (Canceled)

64. (Previously Presented) An isolated nucleic acid molecule encoding an odorant receptor protein comprising seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, wherein the nucleic acid molecule comprises:

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- (a) a nucleic acid sequence given in any one of Figures 9 to 18 (SEQ ID NOs.: 1-10); or
- (b) a nucleic acid sequence degenerate to a sequence of (a) as a result of the genetic code.
- 65. (Previously Presented) An isolated nucleic acid molecule encoding an odorant receptor protein comprising seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, wherein the nucleic acid molecule comprises:
  - (a) a nucleic acid sequence given in any one of the Figures 19 to 31 (SEQ ID NOs.: 11, 13, 15, 17, 19, 21, 23, 25, 27, 29, 31, 33, or 35); or
  - (b) a nucleic acid sequence degenerate to a sequence of(a) as a result of the genetic code.
- 66. (New) A purified odorant receptor protein, wherein the receptor protein comprises seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, and is further characterized by at least one of the following characteristics:
  - (a) the loop between the first transmembrane domain and the second transmembrane domain, and the second transmembrane domain together comprise consecutive amino acids having

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the following sequence: -L, X, X, P, M, Y, X, F, L- (SEQ ID NO: 55);

(b) the third transmembrane domain, and the loop between the third transmembrane domain and the fourth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-M, X, Y, D, R, X, X, A, I, C- (SEQ ID NO: 57); or

-D, R, X, X, A, I, C- (SEQ ID NO: 59);

(c) the loop between the fifth transmembrane domain and the sixth transmembrane domain, and the sixth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-K or R, X, F, S, T, C, X, S, H- (SEQ ID NO: 61); or

-F, S, T, C, X, S, H- (SEQ ID NO: 63); or

(d) the seventh transmembrane domain and the C-terminal domain together comprise consecutive amino acids having one of the following sequences:

-P, X, X, N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 65); or

-P, X, X, N, P, X, I, Y- (SEQ ID NO: 67); or

-N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 69);

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wherein X is any amino acid.

- 67. (New) A purified odorant receptor protein comprising seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains, wherein the protein is selected from the group consisting of:
  - (a) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with tyrosine at position 333 as set forth in row F3 of Figures 4A to 4M (SEQ ID NO: 71),
  - (b) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with glutamine at position 313 as set forth in row F5 of Figures 4A to 4L (SEQ ID NO: 72),
  - (c) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with lysine at position 311 as set forth in row F6 of Figures 4A to 4L (SEQ ID NO: 73),
  - (d) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with glycine at position 317 as set forth in row F12 of Figures 4A to 4L

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(SEQ ID NO: 74),

- (e) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 310 as set forth in row I3 of Figures 4A to 4L (SEQ ID NO: 75),
- (f) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with glycine at position 327 as set forth in row I7 of Figures 4A to 4L (SEQ ID NO: 76),
- (g) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with tryptophan at position 312 as set forth in row I8 of Figures 4A to 4L (SEQ ID NO: 77),
- (h) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 314 as set forth in row I9 of Figures 4A to 4L (SEQ ID NO: 78),
- (i) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 312 as set forth in row I14 of Figures 4A to 4L

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(SEQ ID NO: 79),

- (j) an odorant receptor protein comprising consecutive amino acids having a sequence identical to that beginning with methionine at position 1 and ending with leucine at position 314 as set forth in row I15 of Figures 4A to 4L (SEQ. ID. NO. 80); and
- (k) an odorant receptor protein that shares from 40-80% amino acid identity with any one of the proteins of (a)-(j), comprises seven transmembrane domains, and is further characterized by at least one of the following characteristics:
  - (i) the loop between the first transmembrane domain and the second transmembrane domain, and the second transmembrane domain together comprise consecutive amino acids having the following sequence: -L, X, X, P, M, Y, X, F, L- (SEQ ID NO: 55);
  - (ii) the third transmembrane domain, and the loop between the third transmembrane domain and the fourth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-M, X, Y, D, R, X, X, A, I, C- (SEQ ID NO: 57); or

-D, R, X, X, A, I, C- (SEQ ID NO: 59);

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(iii) the loop between the fifth transmembrane domain and the sixth transmembrane domain, and the sixth transmembrane domain together comprise consecutive amino acids having one of the following sequences:

-K or R, X, F, S, T, C, X, S, H- (SEQ ID NO: 61); or

-F, S, T, C, X, S, H- (SEQ ID NO: 63); or

(iv) the seventh transmembrane domain and the C-terminal domain together comprise consecutive amino acids having one of the following sequences:

-P, X, X, N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 65); or

-P, X, X, N, P, X, I, Y- (SEQ ID NO: 67); or

-N, P, X, I, Y, X, L, R, N- (SEQ ID NO: 69); wherein X is any amino acid.

68. (New) A purified odorant receptor protein comprising seven transmembrane domains and a 17-amino acid cytoplasmic loop between the fifth and sixth transmembrane domains and encoded by a nucleic acid molecule, wherein the nucleic acid molecule comprises a nucleic acid sequence which can be amplified by polymerase chain reaction using:

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- (a) any one of 5' primers A1 (SEQ ID NO: 37), A2 (SEQ ID NO: 38), A3 (SEQ ID NO: 39), A4 (SEQ ID NO: 40), or A5 (SEQ ID NO: 41); and
- (b) any one of 3' primers B1 (SEQ ID NO: 42), B2 (SEQ ID NO: 43), B3 (SEQ ID NO: 44), B4 (SEQ ID NO: 45), B5 (SEQ ID NO: 46), or B6 (SEQ ID NO: 47).